CLAIM AMENDMENTS

- 1. (currently amended) An isolated genomic polynucleotidenucleic acid molecule, said polynucleotide nucleic acid molecule obtainable from human chromosome 7 having a nucleotide sequence at least 95% identical to a sequence selected from the group consisting of:
- (a) a polynucleotide <u>nucleic acid molecule</u> encoding a polypeptide selected from the group consisting of human SNARE YKT6 depicted in SEQ ID NO:1, human liver glucokinase depicted in SEQ ID NO:2, human adipocyte enhancer binding protein <u>1</u> depicted in SEQ ID NO:3 and DNA directed 50kD regulatory subunit (POLD2) depicted in SEQ ID NO:4 and variants thereof;

(c)a polynucleotide nucleic acid molecule selected from the group consisting of SEQ ID NO:5 which encodes human SNARE YKT6 depicted in SEQ ID NO:1, SEQ ID NO:6 which encodes human liver glucokinase depicted in SEQ ID NO:2, SEQ ID NO:8 which encodes human adipocyte enhancer binding protein 1_depicted in SEQ ID NO:3 and SEQ ID NO:7 which encodes DNA directed 50kD regulatory subunit (POLD2) depicted in SEQ ID NO:4 and variants thereof;

- (c) a nucleic acid molecule extending from the 5'-end of SEQ ID NO:5 to the 3'-end of SEQ ID NO:8 that comprises the contiguous coding sequences for SNARE YKT6, glucokinase, POLD2 and the adipocyte enhancer binding protein 1;
 - (c) a polynucleotide which is a variant of SEQ ID NOS:5, 6, 7, or 8;
 - (d) a polynucleotide which is an allelic variant of SEO ID NOS:5, 6, 7, or 8;
 - (e) a polynucleotide which encodes a variant of SEQ ID NOS:1,2, 3, or 4;
- (fd) a polynucleotide nucleic acid molecule which hybridizes to any one of the polynucleotides specified in (a)-(ec)
 - (e) a polynucleotide nucleic acid molecule which is a reverse complement of the polynucleotides specified in (a)- $(f_{\underline{c}})$;
- 2. (currently amended) A nucleic acid construct comprising the polynucleotide nucleic acid molecule of claim 1.
- 3. (currently amended) An expression vector comprising the polynucleotide <u>nucleic</u> acid molecule of claim 1.
- 4. (original) A recombinant host cell comprising the nucleic acid construct molecule of claim 12.

Claim 5 (cancelled)

- 6. (currently amended) A method for obtaining a polypeptide encoded by a polynucleotide nucleic acid molecule obtainable from human chromosome 7, said polypeptide selected from the group consisting of human SNARE YKT6, human liver glucokinase, human adipocyte enhancer binding protein 1 and DNA directed 50kD regulatory subunit (POLD2) comprising:
- (a) culturing the recombinant host cell of claim-54 under conditions that provide for the expression of said polypeptide and
 - (b) recovering said expressed polypeptide.
- 7. (currently amended) A method for preparing an antibody specific to a polypeptide selected from the group consisting of human SNARE YKT6, human liver-glucokinase, human adipocyte enhancer binding protein <u>1</u> and DNA directed 50kD regulatory subunit (POLD2) comprising:
 - (a) obtaining a polypeptide according to the method of claim 6;
 - (b) optionally conjugating said polypeptide to a carrier protein;
- (c) immunizing a host animal with said polypeptide or polypeptide-carrier protein conjugate of step (b) with an adjuvant and
 - (d) obtaining antibody from said immunized host animal.
- 8. (currently amended) An antisense oligonucleotide or mimetic to an isolated polynucleotide isolated nucleic acid molecule of at least 15 nucleotides or mimetic which hybridizes at high stringency to a non-coding region of specific to SEQ ID NOS:5, 6, 7 or 8the nucleic acid molecule of claim 1, which non-coding region is selected from the group consisting of an intron, a splice junction, a 5' non-coding region, a transcription factor binding region, an expression control region and a 3' non-coding region.
- 9. (currently amended) A method of diagnosing a pathological condition or susceptibility to a pathological condition in a subject comprising:

- (b) diagnosing a pathological condition or a susceptibility to a pathological condition based on the presence or absence of said mutation
 - (a) <u>isolating genomic DNA from a subject;</u>
 - (b) determining the presence or absence of a variant in said genomic DNA using the nucleic acid molecule of claim 8 and
 - (c) diagnosing a pathological condition or a susceptibility to a pathological condition based on the presence or absence of said variant.
- 10. (currently amended) A composition comprising the polynucleotide nucleic acid molecule of claim 1 and a carrier.
- 11. (currently amended) A composition comprising the antisense oligonucleotidenucleic acid molecule of claim 8 and a carrier.
- 12. (original) A method for preventing, treating or ameliorating a medical condition, comprising administering to a subject an amount of the composition of claim 10 effective to prevent, treat or ameliorate said medical condition.
- 13. (original) A method for preventing, treating or ameliorating a medical condition, comprising administering to a subject an amount of the composition of claim 11 effective to prevent, treat or ameliorate said medical condition.
- 14. (currently amended) A kit comprising the polynucleotide nucleic acid molecule of claim 18.
- 15. (original) The kit according to claim 14, in which the polynucleotide is labeled with a detectable substance.
- 16. (currently amended) A kit comprising the antisense oligonucleotide or mimetic of claim 8. The kit according to claim 14, which comprises a plurality of nucleic acid molecules.

- 23. (new) A method for modulating levels of human SNARE YKT6, human glucokinase, human adipocyte enhancer binding protein 1 or DNA directed 50kD regulatory subunit (POLD2) in a subject in need thereof comprising administering to said subject an amount of the nucleic acid molecule of claim 1 effective to modulate said human SNARE YKT6, human glucokinase, human adipocyte enhancer binding protein 1 or DNA directed 50kD regulatory subunit (POLD2) levels.
- 24. (new) A method for modulating levels of human SNARE YKT6, human glucokinase, human adipocyte enhancer binding protein 1 or DNA directed 50kD regulatory subunit (POLD2) in a subject in need thereof comprising administering to said subject an amount of the nucleic acid molecule of claim 8 effective to modulate said human SNARE YKT6, human glucokinase, human adipocyte enhancer binding protein 1 or DNA directed 50kD regulatory subunit (POLD2) levels.
- 25. (new) A method of identifying variants of SEQ ID NOS: 5, 6, 7 or 8 comprising
- (a) isolating genomic DNA from a subject and
- (b) determining the presence or absence of a variant in said genomic DNA using the nucleic acid molecule of claim 8.
- 26. (new) A method for detecting the presence or absence of a non-coding nucleic acid sequence specific to the nucleic acid molecule of claim 1 in a sample, said method comprising contacting the sample with a nucleic acid molecule of at least 15 nucleotides which hybridizes at high stringency to a non-coding region specific to the nucleic acid molecule of claim 1, which non-coding region is selected from the group consisting of an intron, a splice junction, a 5' non-coding region, a transcription factor binding region, an expression control region and a 3' non-coding region.